

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 09/486,247B
Source: IFW/6
Date Processed by STIC: 5/20/05

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 05/20/2005

PATENT APPLICATION: US/09/486,247B

TIME: 09:07:27

Input Set : A:\31304-703.831.ST25.txt

Output Set: N:\CRF4\05202005\I486247B.raw

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3 <110> APPLICANT: Deutsches Krebsforschungszentrum Stiftung des öffentlichen
4   Rechts
5   DEAR, Terence N.
6   BOEHM, Thomas
8 <120> TITLE OF INVENTION: PROTEASE-RELATED PROTEIN
10 <130> FILE REFERENCE: 31304-703.831
12 <140> CURRENT APPLICATION NUMBER: 09/486,247B
13 <141> CURRENT FILING DATE: 2000-05-25
15 <160> NUMBER OF SEQ ID NOS: 8
17 <170> SOFTWARE: PatentIn version 3.2
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 822
21 <212> TYPE: DNA
22 <213> ORGANISM: Mouse
24 <400> SEQUENCE: 1
25 taggtggtgt cattccccctc caacctgagt gctggcaggt acactgctgg ccaccagcag      60
27 atgcccataga agatgctgac aatgaagatg ctggccctgt gcttggttct tgctaaatca      120
29 gcctggtcgg aggaacagga gaaggtggtt catggaggcc cgtgtttgaa ggactccac      180
31 cctttccagg ctgccctcta cacctcaggt cacttgctgt gtggtggggt cctcattgac      240
33 ccacagtggg tgctgacagc tgcccactgc aaaaaaccga atctgcaggt gatcttgggg      300
35 aaacacaacc tacggcaaac agagactttc caaaggcaaa tctcagtga caggactatt      360
37 gtccatcccc gctacaacct tgaaaccac gacaatgaca tcatgatggt gcatctgaaa      420
39 aatccagtca aattctctaa aaagatccag cctctgccct tgaagaatga ctgctctgag      480
41 gagaatccca actgccagat cctgggctgg ggcaagatgg aaaatggtga cttcccagat      540
43 accattcagt gtgctgatgt ccatctggtg ccccgaggagc agtgtgagcg tgcctaccct      600
45 ggcaagatca cccagagcat ggtgtgcgca ggcgacatga aagaaggcaa cgattcctgt      660
47 cagggtgatt ctggaggtcc cctagtatgt gggggtcgcc tccgagggct cgtgtcatgg      720
49 ggtgacatgc cctgtggatc aaaggagaag ccaggagttt acaccgatgt ctgcactcat      780
51 atcagatgga tccaaaacat cctcagaaac aagtggctgt ga      822
54 <210> SEQ ID NO: 2
55 <211> LENGTH: 253
56 <212> TYPE: PRT
57 <213> ORGANISM: Mouse
59 <400> SEQUENCE: 2
61 Met Pro Met Lys Met Leu Thr Met Lys Met Leu Ala Leu Cys Leu Val
62 1          5          10          15
65 Leu Ala Lys Ser Ala Trp Ser Glu Glu Gln Glu Lys Val Val His Gly
66          20          25          30
69 Gly Pro Cys Leu Lys Asp Ser His Pro Phe Gln Ala Ala Leu Tyr Thr
70          35          40          45
73 Ser Gly His Leu Leu Cys Gly Gly Val Leu Ile Asp Pro Gln Trp Val
74          50          55          60
77 Leu Thr Ala Ala His Cys Lys Lys Pro Asn Leu Gln Val Ile Leu Gly

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78 65          70          75          80
81 Lys His Asn Leu Arg Gln Thr Glu Thr Phe Gln Arg Gln Ile Ser Val
82          85          90          95
85 Asp Arg Thr Ile Val His Pro Arg Tyr Asn Pro Glu Thr His Asp Asn
86          100          105          110
89 Asp Ile Met Met Val His Leu Lys Asn Pro Val Lys Phe Ser Lys Lys
90          115          120          125
93 Ile Gln Pro Leu Pro Leu Lys Asn Asp Cys Ser Glu Glu Asn Pro Asn
94          130          135          140
97 Cys Gln Ile Leu Gly Trp Gly Lys Met Glu Asn Gly Asp Phe Pro Asp
98 145          150          155          160
101 Thr Ile Gln Cys Ala Asp Val His Leu Val Pro Arg Glu Gln Cys Glu
102          165          170          175
105 Arg Ala Tyr Pro Gly Lys Ile Thr Gln Ser Met Val Cys Ala Gly Asp
106          180          185          190
109 Met Lys Glu Gly Asn Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu
110          195          200          205
113 Val Cys Gly Gly Arg Leu Arg Gly Leu Val Ser Trp Gly Asp Met Pro
114          210          215          220
117 Cys Gly Ser Lys Glu Lys Pro Gly Val Tyr Thr Asp Val Cys Thr His
118 225          230          235          240
121 Ile Arg Trp Ile Gln Asn Ile Leu Arg Asn Lys Trp Leu
122          245          250

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125 <210> SEQ ID NO: 3

126 <211> LENGTH: 12

127 <212> TYPE: DNA

128 <213> ORGANISM: Artificial

130 <220> FEATURE:

131 <223> OTHER INFORMATION: Oligonucleotide adaptor forrepresentational difference

analysis

133 <400> SEQUENCE: 3

134 gatctgcggt ga 12

137 <210> SEQ ID NO: 4

138 <211> LENGTH: 24

139 <212> TYPE: DNA

140 <213> ORGANISM: Artificial

142 <220> FEATURE:

143 <223> OTHER INFORMATION: Oligonucleotide adaptor forrepresentational difference

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145 <400> SEQUENCE: 4

146 agcactctcc agcctctcac cgca 24

149 <210> SEQ ID NO: 5.

150 <211> LENGTH: 12

151 <212> TYPE: DNA

152 <213> ORGANISM: Artificial

154 <220> FEATURE:

155 <223> OTHER INFORMATION: Oligonucleotide adaptor forrepresentational difference

analysis

157 <400> SEQUENCE: 5

158 gatctgttca tg 12

161 <210> SEQ ID NO: 6

162 <211> LENGTH: 24

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163 <212> TYPE: DNA
164 <213> ORGANISM: Artificial
166 <220> FEATURE:
167 <223> OTHER INFORMATION: Oligonucleotide adaptor forrepresentational difference
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169 <400> SEQUENCE: 6
170 accgacgtcg actatccatg aaca 24
173 <210> SEQ ID NO: 7
174 <211> LENGTH: 12
175 <212> TYPE: DNA
176 <213> ORGANISM: Artificial
178 <220> FEATURE:
179 <223> OTHER INFORMATION: Oligonucleotide adaptor forrepresentational difference
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181 <400> SEQUENCE: 7
182 gatcttccct cg 12
185 <210> SEQ ID NO: 8
186 <211> LENGTH: 24
187 <212> TYPE: DNA
188 <213> ORGANISM: Artificial
190 <220> FEATURE:
191 <223> OTHER INFORMATION: Oligonucleotide adaptor forrepresentational difference
analysis
193 <400> SEQUENCE: 8
194 aggcaactgt gctatccgag ggaa 24

```

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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:3,4,5,6,7,8

VERIFICATION SUMMARY

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